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==============================================================
Sequence Listing was accepted.
See attached Validation Report.
If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).
Reviewer: Durreshwar Anjum
Timestamp: [year=2009; month=4; day=27; hr=13; min=29; sec=4; ms=718; ]
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Application No: 10567074   Version No: 2.0

                    Input Set:

                    Output Set:

                    Started: 2009-04-14 17:39:59.103
                    Finished: 2009-04-14 17:40:03.182
                    Elapsed: 0 hr(s) 0 min(s) 4 sec(s) 79 ms
            Total Warnings: 29
              Total Errors: 1
    No. of SeqIDs Defined: 60
        Actual SeqID Count: 60

Error code   Error Description

W 402   Undefined organism found in <213> in SEQ ID (3)

E 201   Mandatory field data missing in <223> in SEQ ID (3)

W 402   Undefined organism found in <213> in SEQ ID (4)

W 402   Undefined organism found in <213> in SEQ ID (5)

W 402   Undefined organism found in <213> in SEQ ID (16)

W 402   Undefined organism found in <213> in SEQ ID (18)

W 402   Undefined organism found in <213> in SEQ ID (29)

W 402   Undefined organism found in <213> in SEQ ID (30)

W 402   Undefined organism found in <213> in SEQ ID (31)

W 402   Undefined organism found in <213> in SEQ ID (32)

W 402   Undefined organism found in <213> in SEQ ID (33)

W 402   Undefined organism found in <213> in SEQ ID (34)

W 402   Undefined organism found in <213> in SEQ ID (35)

W 402   Undefined organism found in <213> in SEQ ID (38)

W 402   Undefined organism found in <213> in SEQ ID (41)

W 402   Undefined organism found in <213> in SEQ ID (42)

W 402   Undefined organism found in <213> in SEQ ID (44)

W 402   Undefined organism found in <213> in SEQ ID (47)

W 402   Undefined organism found in <213> in SEQ ID (49)

W 402   Undefined organism found in <213> in SEQ ID (50)

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          Input Set:


          Output Set:



            Started:   2009-04-14 17:39:59.103
           Finished:   2009-04-14 17:40:03.182
            Elapsed:   0 hr(s) 0 min(s) 4 sec(s) 79 ms
     Total Warnings:   29
       Total Errors:   1
No. of SeqIDs Defined:   60
  Actual SeqID Count:    60
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Error code	Error Description
W 402	Undefined organism found in <213> in SEQ ID (51) This error has occured more than 20 times, will not be displayed
W 213	Artificial or Unknown found in <213> in SEQ ID (53)
W 213	Artificial or Unknown found in <213> in SEQ ID (54)
W 213	Artificial or Unknown found in <213> in SEQ ID (55)
W 213	Artificial or Unknown found in <213> in SEQ ID (56)
W 213	Artificial or Unknown found in <213> in SEQ ID (57)
W 213	Artificial or Unknown found in <213> in SEQ ID (58)
W 213	Artificial or Unknown found in <213> in SEQ ID (59)
W 213	Artificial or Unknown found in <213> in SEQ ID (60)

<110> Scherer, Stephen W.

   Minassian, Berge A.

<120> Lafora's Disease Gene

<160> 60

<170> PatentIn version 3.1

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ccggccgccc atcgcgccgc ccccagcgcc cccggagccc tcacctgcca ccacaccttc      360

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caccccctgg ccctggggac tggggtttgc agcaccaggg tgaaagtgtt tagctcaagt      900

atgcagcttg tcggccaagt ggataccttt gggctgagcc tctactttcc ctccaaaata      960

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gagaaaaagg attttgtggt ggccatcact ttcattcaat ttgcgactgc tctgatggca   2040

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Phe Gly His Arg Gln Gln Arg Arg Pro Arg Asn Leu Ser Cys Gly His
        35   40   45

Val Val Cys Leu Ala Cys Val Ala Ala Leu Ala His Pro Arg Thr Leu
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Ala Leu Glu Cys Pro Phe Cys Arg Arg Ala Cys Arg Gly Cys Asp Thr
65   70   75   80

Ser Asp Cys Leu Pro Val Leu His Leu Ile Glu Leu Leu Gly Ser Ala
                85   90   95

Leu Arg Gln Ser Pro Ala Ala His Arg Ala Ala Pro Ser Ala Pro Gly
            100   105   110

Ala Leu Thr Cys His His Thr Phe Gly Gly Trp Gly Thr Leu Val Asn
        115   120   125

Pro Thr Gly Leu Ala Leu Cys Pro Lys Thr Gly Arg Val Val Val Val
    130   135   140

His Asp Gly Arg Arg Arg Val Lys Ile Phe Asp Ser Gly Gly Gly Cys
145   150   155   160

Ala His Gln Phe Gly Glu Lys Gly Asp Ala Ala Gln Asp Ile Arg Tyr
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Pro Val Asp Val Thr Ile Thr Asn Asp Cys His Val Val Val Thr Asp
            180   185   190

Ala Gly Asp Arg Ser Ile Lys Val Phe Asp Phe Phe Gly Gln Ile Lys
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Leu Val Ile Gly Gly Gln Phe Ser Leu Pro Trp Gly Val Glu Thr Thr
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Pro Gln Asn Gly Ile Val Val Thr Asp Ala Glu Ala Gly Ser Leu His
225   230   235   240

Leu Leu Asp Val Asp Phe Ala Glu Gly Val Leu Arg Arg Thr Glu Arg
            245   250   255

Leu Gln Ala His Leu Cys Asn Pro Arg Gly Val Ala Val Ser Trp Leu
            260   265   270

Thr Gly Ala Ile Ala Val Leu Glu His Pro Leu Ala Leu Gly Thr Gly
            275   280   285

Val Cys Ser Thr Arg Val Lys Val Phe Ser Ser Ser Met Gln Leu Val
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Gly Gln Val Asp Thr Phe Gly Leu Ser Leu Tyr Phe Pro Ser Lys Ile
305   310   315   320

Thr Ala Ser Ala Val Thr Phe Asp His Gln Gly Asn Val Ile Val Ala
            325   330   335

Asp Thr Ser Gly Pro Ala Ile Leu Cys Leu Gly Lys Pro Glu Glu Phe
            340   345   350

Pro Val Pro Lys Pro Met Val Thr His Gly Leu Ser His Pro Val Ala
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Leu Thr Phe Thr Lys Glu Asn Ser Leu Leu Val Leu Asp Thr Ala Ser
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His Ser Ile Lys Val Tyr Lys Val Asp Trp Gly
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ccgccggggg accgcaggcc gcggccgaga ggctgcgcgc tgcgcccgcg acgtcaggcc     600

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accgagcggc gcccgcggga gcggcggcgg ccgcgcg atg ggg gcc gaa gcg gcg       715
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ggg agc ggg cgg gcg ctg cgg gag ctg gtg cgc gag gcc gag gtc agc       763
Gly Ser Gly Arg Ala Leu Arg Glu Leu Val Arg Glu Ala Glu Val Ser
         10                  15                  20

ttg ctc gag tgc aag gtg tgc ttc gag agg ttc ggc cac cgc cag cag       811
Leu Leu Glu Cys Lys Val Cys Phe Glu Arg Phe Gly His Arg Gln Gln
     25                  30                  35

cgg cgc ccg cgc aac ctg ccc tgc ggc cac gtg gtg tgc ctg gcc tgc       859
Arg Arg Pro Arg Asn Leu Pro Cys Gly His Val Val Cys Leu Ala Cys
     40                  45                  50

gtg gcg gcc ctg gcg cac ccg cgg acg ctg gcc ctg gag tgc ccc ttc       907
Val Ala Ala Leu Ala His Pro Arg Thr Leu Ala Leu Glu Cys Pro Phe
55                  60                  65                  70
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tgc cgc cgg gcc tgc cgc ggc tgc gac acc agc gac tgc ctg ccg gtg          955
Cys Arg Arg Ala Cys Arg Gly Cys Asp Thr Ser Asp Cys Leu Pro Val
                75                  80                  85


ctt cac ctc ctg gag ctc ctg ggc tcg gcg ctg cgc cca gcc ccc gcc         1003
Leu His Leu Leu Glu Leu Leu Gly Ser Ala Leu Arg Pro Ala Pro Ala
                90                  95                  100


gcc ccc cgc gcc gcc ccc cgc gcc gcc ccc tgc gcc ccg ggc gcc ctc         1051
Ala Pro Arg Ala Ala Pro Arg Ala Ala Pro Cys Ala Pro Gly Ala Leu
        105                 110                 115


gcc tgc cat cac gcg ttc gga ggc tgg ggg acc ctg gtc aac ccc acg         1099
Ala Cys His His Ala Phe Gly Gly Trp Gly Thr Leu Val Asn Pro Thr
    120                 125                 130


ggg ctg gcg ctg tgc ccc aag acc ggg cgg gtc gtg gtg gtg cac gac         1147
Gly Leu Ala Leu Cys Pro Lys Thr Gly Arg Val Val Val Val His Asp
135                 140                 145                 150


ggc agg agg cgg gtc aag atc ttt gac tcc ggg gga gga tgc gcc cat         1195
Gly Arg Arg Arg Val Lys Ile Phe Asp Ser Gly Gly Gly Cys Ala His
                155                 160                 165


cag ttt gga gag aag ggg gag gct gcc cag gac att agg tac ccc ctg         1243
Gln Phe Gly Glu Lys Gly Glu Ala Ala Gln Asp Ile Arg Tyr Pro Leu
                170                 175                 180


gac gtc gcc gtc acc aac gac tgc cac gtg gtt gtc acc gac gcc ggc         1291
Asp Val Ala Val Thr Asn Asp Cys His Val Val Val Thr Asp Ala Gly
        185                 190                 195


gac cgc tcc atc aaa gtg ttt gat ttc ttt ggc cag atc aag ctc gtc         1339
Asp Arg Ser Ile Lys Val Phe Asp Phe Phe Gly Gln Ile Lys Leu Val
    200                 205                 210


att gga gac cag ttt tcc tta cct tgg ggc gtg gag acc acc cct cag         1387
Ile Gly Asp Gln Phe Ser Leu Pro Trp Gly Val Glu Thr Thr Pro Gln
215                 220                 225                 230


aat ggg gtc gtg gta act gac gcc gag gca ggg tcg ctg cac ctg ctg         1435
Asn Gly Val Val Val Thr Asp Ala Glu Ala Gly Ser Leu His Leu Leu
                235                 240                 245


gaa gtc gac ttt gca gaa gga gcc ctc cag agg act gaa aag ctg caa         1483
Glu Val Asp Phe Ala Glu Gly Ala Leu Gln Arg Thr Glu Lys Leu Gln
            250                 255                 260


ggt cat ctg tgc aac ccg cga ggg gtg gcc gtg tcc tgg ctc act ggg         1531
Gly His Leu Cys Asn Pro Arg Gly Val Ala Val Ser Trp Leu Thr Gly
        265                 270                 275


gcc att gcg gtc ctg gag cac cct ccg ggg ctg ggg gct ggg gcg ggc         1579
Ala Ile Ala Val Leu Glu His Pro Pro Gly Leu Gly Ala Gly Ala Gly
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Ser Thr Ala Val Lys Val Phe Ser Pro Thr Met Gln Leu Ile Gly Gln
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gtg gat acc ttt ggg ctc agc ctc ttt ttc ccc tct aga ata acc gcc    1675
Val Asp Thr Phe Gly Leu Ser Leu Phe Phe Pro Ser Arg Ile Thr Ala
                315             320             325

tcc gcc gtg acc ttt gat cac cag ggg aat gtg att gtt gca gat act    1723
Ser Ala Val Thr Phe Asp His Gln Gly Asn Val Ile Val Ala Asp Thr
        330             335             340

tct agt cag gcc gtc cta tgc ttg gga cag cct gag gaa ttt cca gtc    1771
Ser Ser Gln Ala Val Leu Cys Leu Gly Gln Pro Glu Glu Phe Pro Val
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ctg aag ccc atc atc acc cat ggt ctt tcc cat cct gtg gca ctg acc    1819
Leu Lys Pro Ile Ile Thr His Gly Leu Ser His Pro Val Ala Leu Thr
    360             365             370

ttc acc aag gag aat tct ctt ctt gtg ctg gac agt gca gcc cat tcc    1867
Phe Thr Lys Glu Asn Ser Leu Leu Val Leu Asp Ser Ala Ala His Ser
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Val Lys Val Tyr Lys Ala Asp Trp Gly
                395

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tgagaaaact gagacccgag ggtttaagtg attctctgaa ggtcatgttt acttactgtg  2217

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            20   25   30

Phe Gly His Arg Gln Gln Arg Arg Pro Arg Asn Leu Pro Cys Gly His
        35   40   45

Val Val Cys Leu Ala Cys Val Ala Ala Leu Ala His Pro Arg Thr Leu
   50   55   60

Ala Leu Glu Cys Pro Phe Cys Arg Arg Ala Cys Arg Gly Cys Asp Thr
65   70   75   80

Ser Asp Cys Leu Pro Val Leu His Leu Leu Glu Leu Leu Gly Ser Ala
                85   90   95

Leu Arg Pro Ala Pro Ala Ala Pro Arg Ala Ala Pro Arg Ala Ala Pro
                100   105   110

Cys Ala Pro Gly Ala Leu Ala Cys His His Ala Phe Gly Gly Trp Gly
          115   120   125